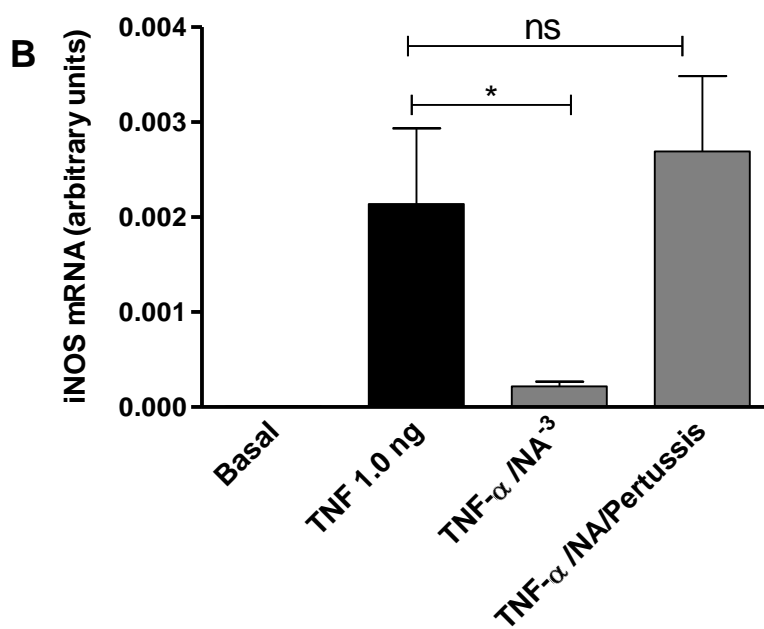
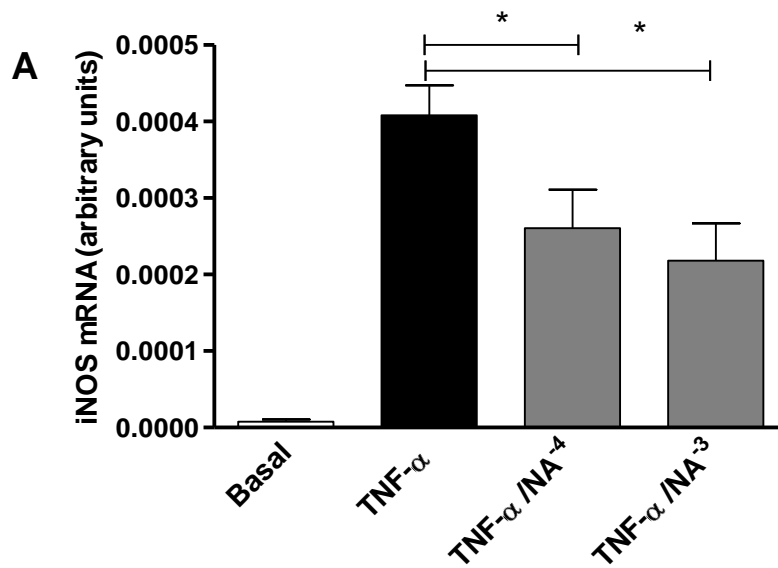
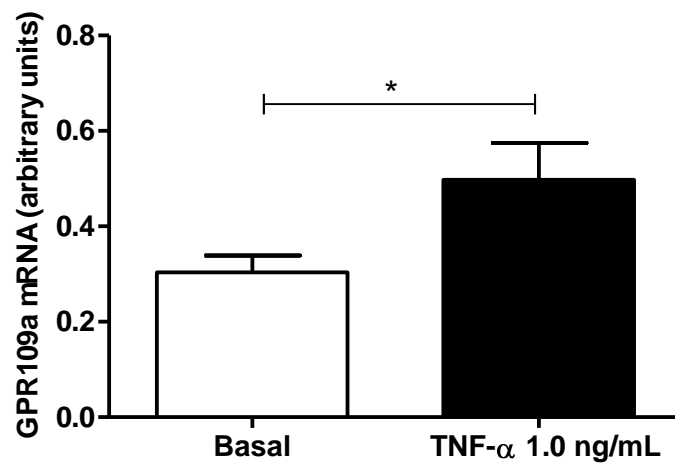


Supplementary  
Figure (i)



Supplementary  
Figure (ii)



Supplementary  
Figure (iii)

## **Supplementary Methods and Results**

### **Cell viability methods**

Cell viability was measured in 3T3-L1 cells differentiated for 10 days then treated with varying doses of TNF- $\alpha$ . Viability was measured using The CellTiter 96<sup>®</sup> AQueous One Solution Cell Proliferation Assay (Promega, Southampton, UK). Absorbance at 490 nm was measured and recorded as mean fluorescence measured by relative light units.

### **Measurement of iNOS and GPR109a Gene expression in 3T3-L1 Adipocytes**

Total RNA was prepared from cell lysates of 3T3-L1 adipocytes differentiated for 10-14 days. Total RNA was purified using Qiagen RNEasy mini columns and 1  $\mu$ g was reverse transcribed using a QuantiTect<sup>®</sup> Reverse Transcription Kit containing a genomic DNA wipe-out step and using Oligo dT's and random hexamers as primers.

Inducible nitric oxide synthase (iNOS) mRNA expression was measured by Taqman<sup>™</sup> Gene Expression assay ID Mm01309902\_m1 according to manufacturer's instructions (supplementary figure (ii)).

GPR 109a mRNA expression was measured by Real-time PCR using Sybr Green Mastermix (Applied Biosystems, Warrington, UK) and sense and anti-sense primers (0.25  $\mu$ M final concentration). Primer sequences were sense; 5'

ATGAAAACATCGCCAAGGTC 3' and antisense; 5'

TGGATTTCCAGGACTTGAGG 3', expected product size, 117 base pairs.

Cycling parameters were as follows: activation of Taq polymerase, 10 min at 95°C, then 40 cycles at 95°C for 15 sec, then extension at 60°C for 1 min, followed by a melt curve analysis. PCR products were sequenced to confirm identity using the Illumina

Genome Analysis system. Sequencing results were subjected to BlastN analysis and confirmed 100% identity with mouse GPR109a. To measure a whether TNF- $\alpha$  was able to affect gene expression of GPR109a, differentiated 3T3-L1 adipocytes were incubated with TNF- $\alpha$  1.0 ng/mL for 4 h and mRNA was measured by quantitative RT-PCR (supplementary figure iii).

**Supplementary Figure (i)** Cell viability in 3T3-L1 cells treated with varying doses of TNF- $\alpha$  was measured using CellTiter 96<sup>®</sup> Aqueous One Solution Cell Proliferation Assay (Promega, Southampton, UK). Absorbance at 490 nm was measured and recorded.

**Supplementary Figure (ii) A.** mRNA levels of iNOS, for RNA analyses, cells were incubated for 4 h and for media analysis cells were incubated for 24 h, with DMEM only (Basal), DMEM + TNF- $\alpha$  1.0 ng / ml (TNF- $\alpha$  ), DMEM + TNF- $\alpha$  1.0 ng / ml + nicotinic acid  $10^{-4}$  M (TNF- $\alpha$  + NA<sup>-4</sup>) or DMEM + TNF- $\alpha$  1.0 ng / ml + nicotinic acid  $10^{-3}$  M (TNF- $\alpha$  + NA<sup>-3</sup>). N = 6 for each treatment, \* $P < 0.05$ , via one-way ANOVA with Bonferroni's multiple comparison post-hoc test.

**B.** The effect of pre-treatment with pertussis (PTX) on adipocyte mRNA expression of iNOS. N=6 for each treatment, \* $P < 0.05$ .

**Supplementary Figure (iii)** TNF- $\alpha$  exposure 1.0 ng/mL for 4h upregulates GPR109a mRNA in 3T3-L1 differentiated adipocytes (\*  $P < 0.05$ , n = 6).